

(79)

CTCGGTGTCCTCTTGCTGTCCAGCTCTGCAGCCTCCGGCGCCCTCCCGCCACGCCATGGACGCCAAGGTCTCGCCC
MetAspAlaLysValValAla

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(241)

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*

(403)

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Lys

GGGAGGAGGGCCTGAACCCCTGGCCAGGATGGCCGGGGGACAGCACTGACTGGGGTTCATGGTAAGGTTTGCCAGCATAAA

GACACTCCGCCATAGCATATGGTACGATATTGCAGCTTATATTATCCCTGCCCTCGCCCGTGACAAATGGAGCTTTTATA

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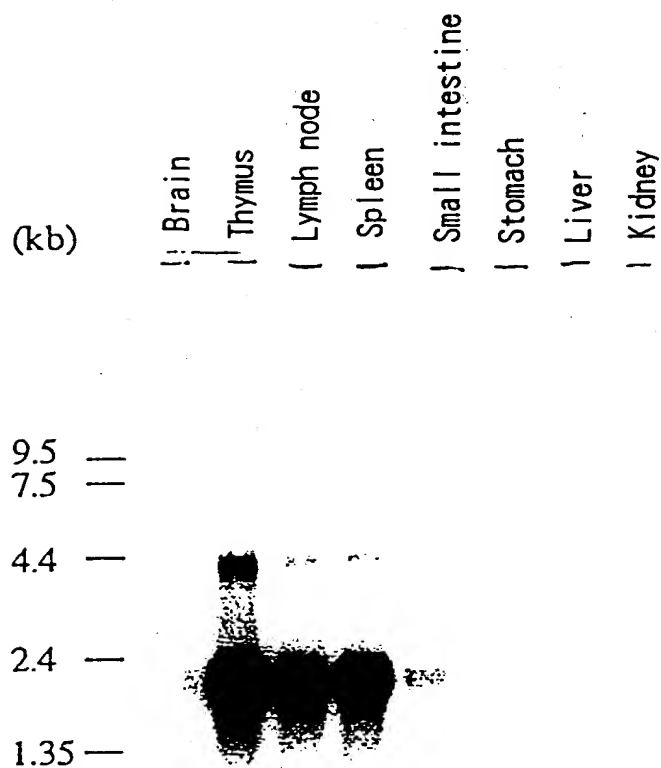
Fig. 1

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GGTCGGGAAGCTTCCCTTTAGAAAGTGATCCCTGAAGCTGTGCTCAGAGACCCCTTTCCCTAGCCGTTCCCTCTCTCTGCTTGC
(1051)
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AGAATGGAGCCAGACCATCCTGGATAATGTGAGAACATGCCCTAGATTTTACCCACAAACACAAAGTCTGAGAAITTAATCATA
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TTCTGGGTTCCAATCAGAAATGGAGACAAGATAAAACTTGCATACATTTCTTACGATCACAGACGGCCCTGGTGGTTTTTGG
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(1699)
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Fig. 1 (Continued)

A



B

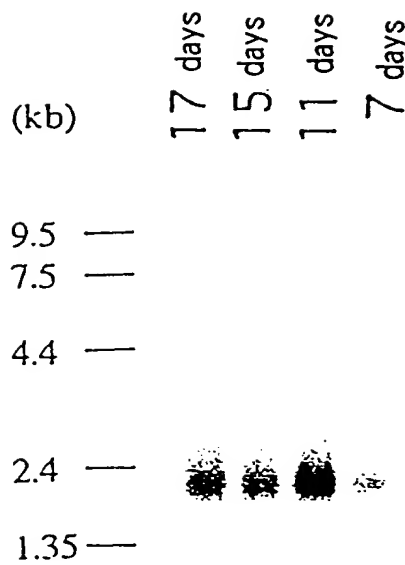


Fig. 2

CHO cells

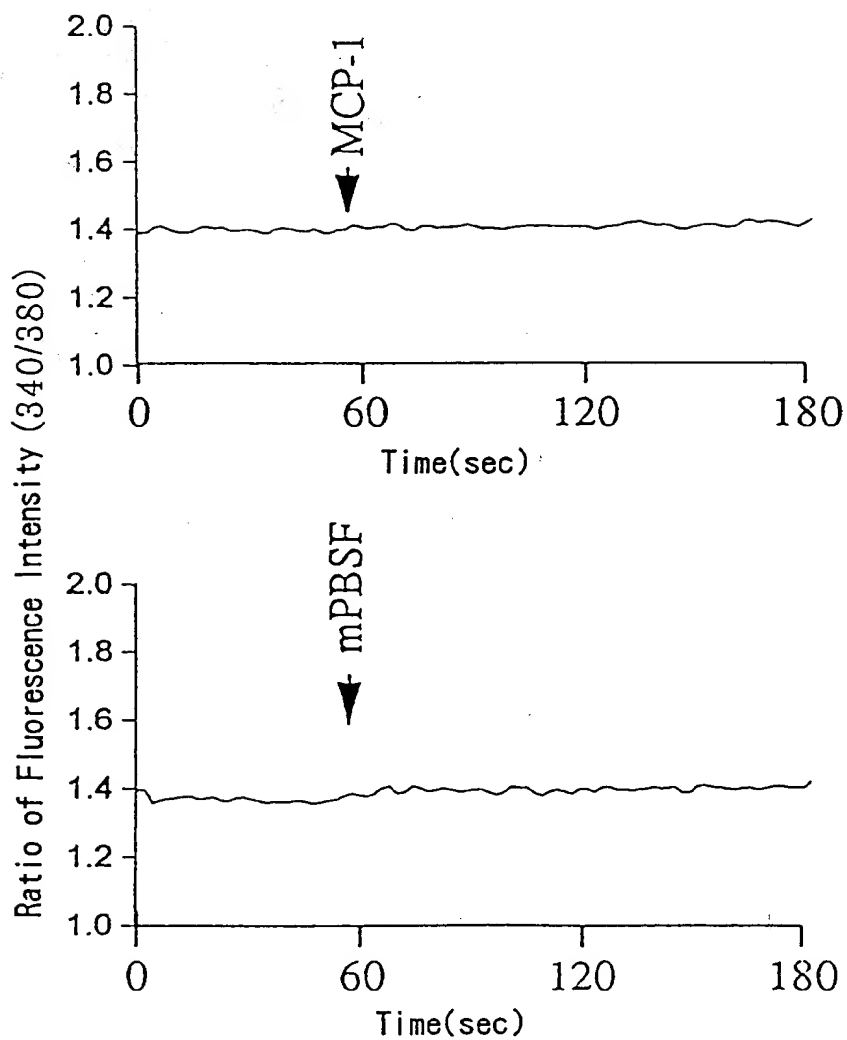


Fig. 3

CHO cells (CC CKR2B+)

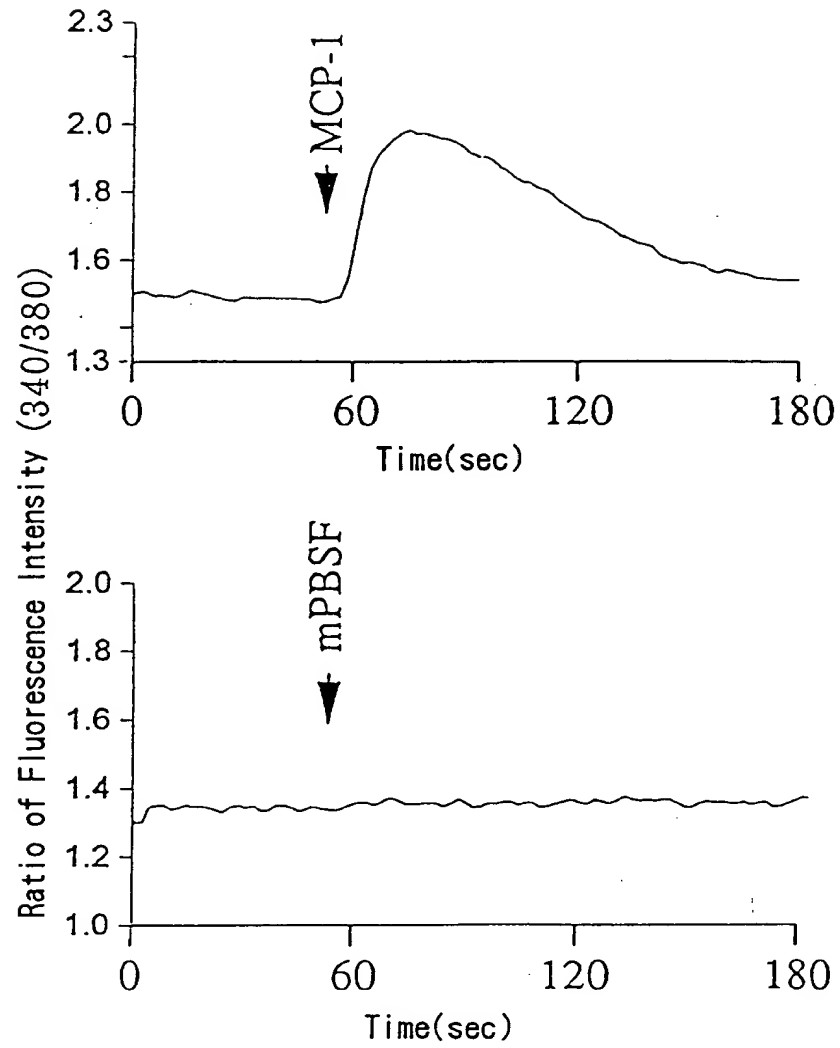


Fig. 4

CHO cells (mPBSF/SDF-1R+)

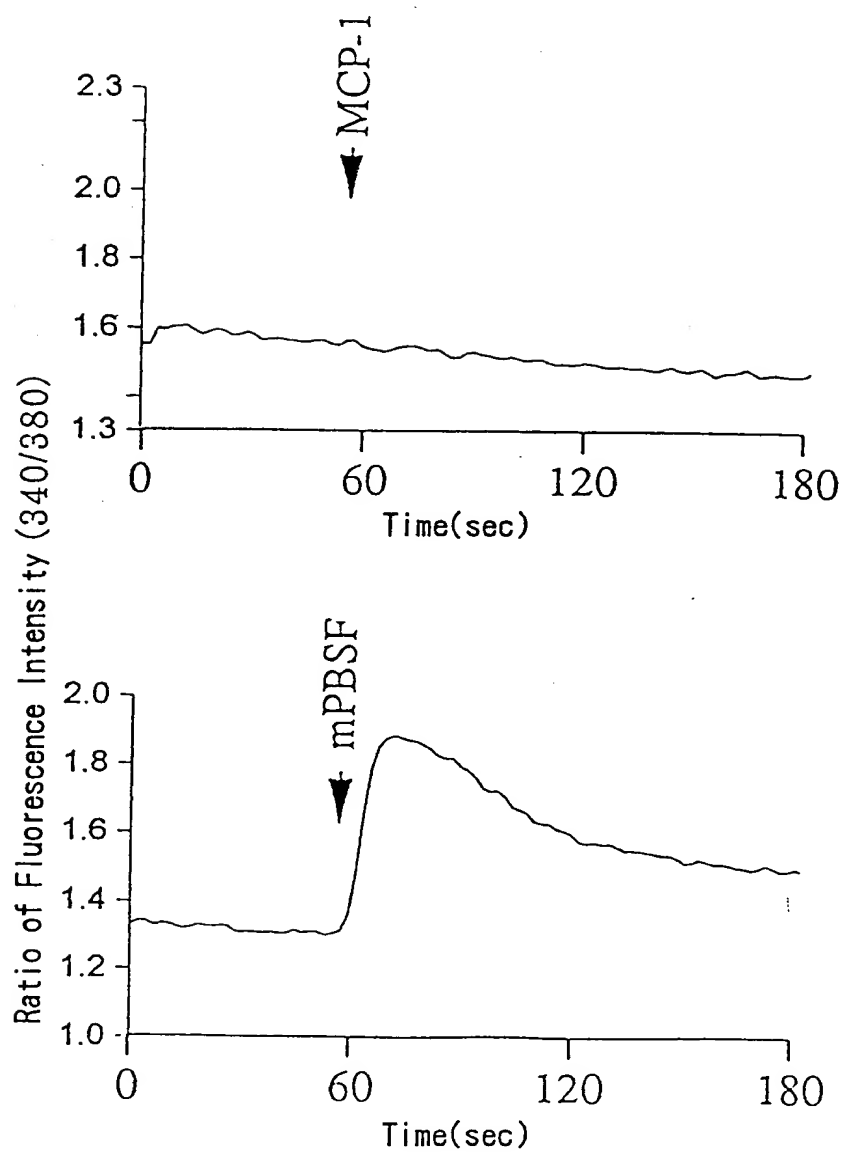


Fig. 5

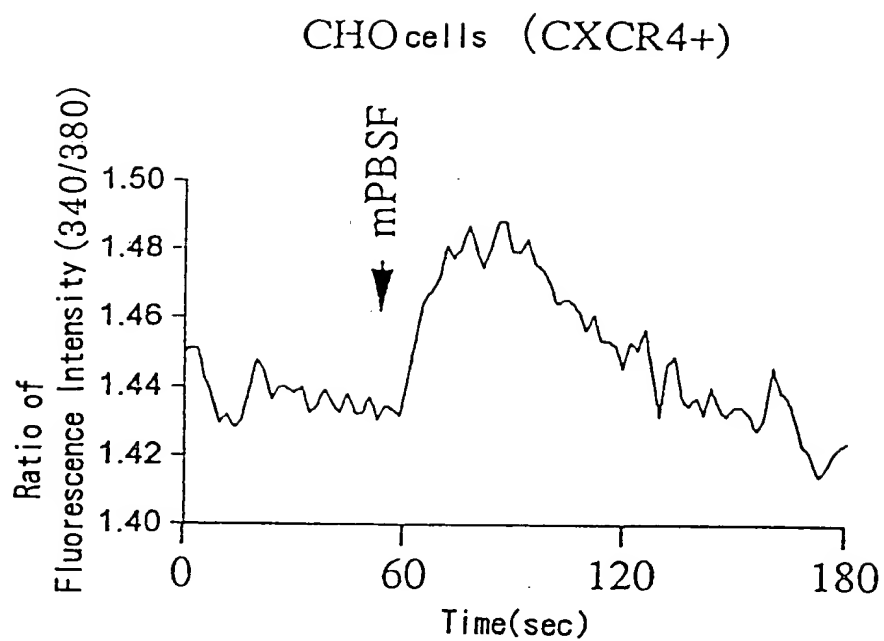


Fig. 6

CHO cells (mPBSF/SDF-1R+)

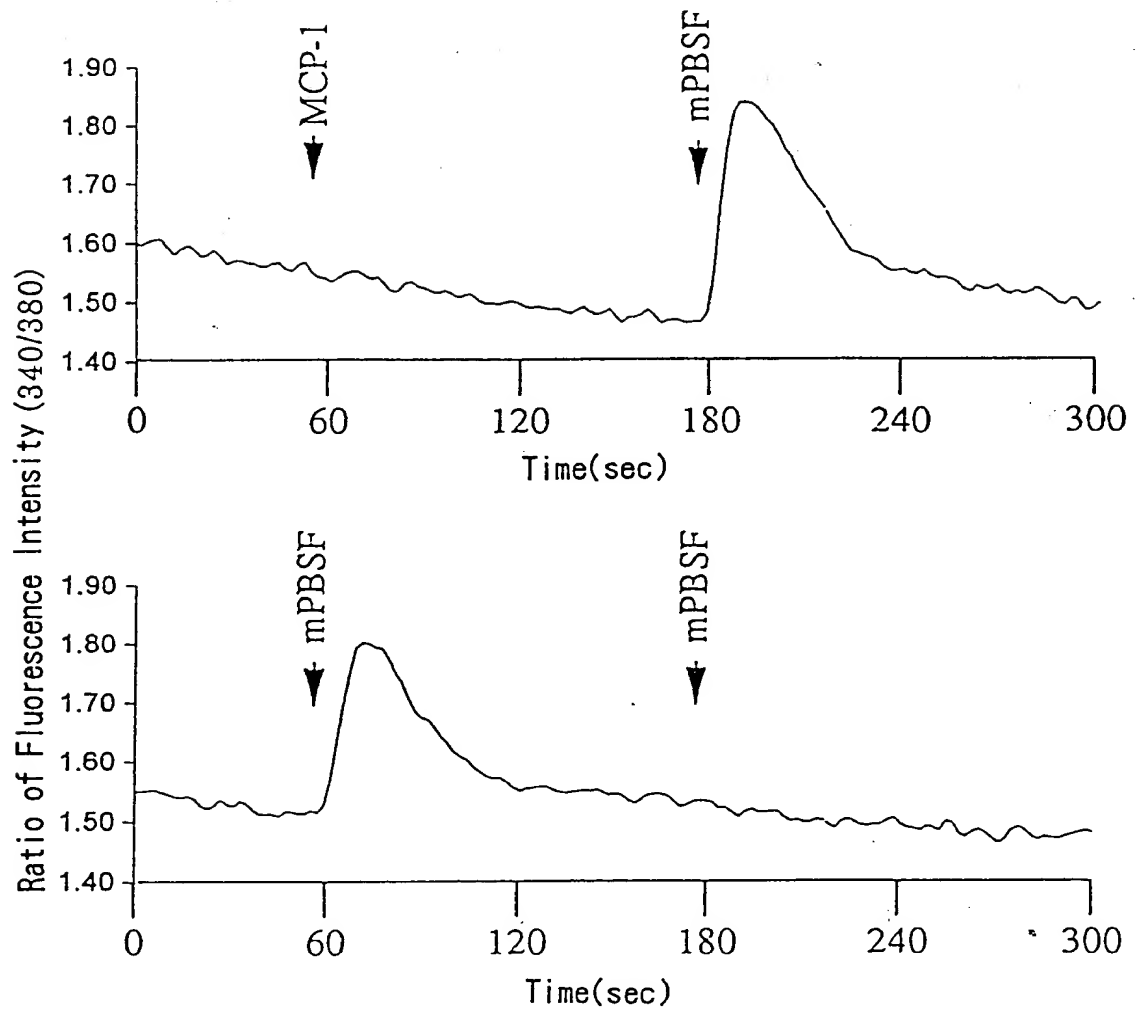


Fig. 7

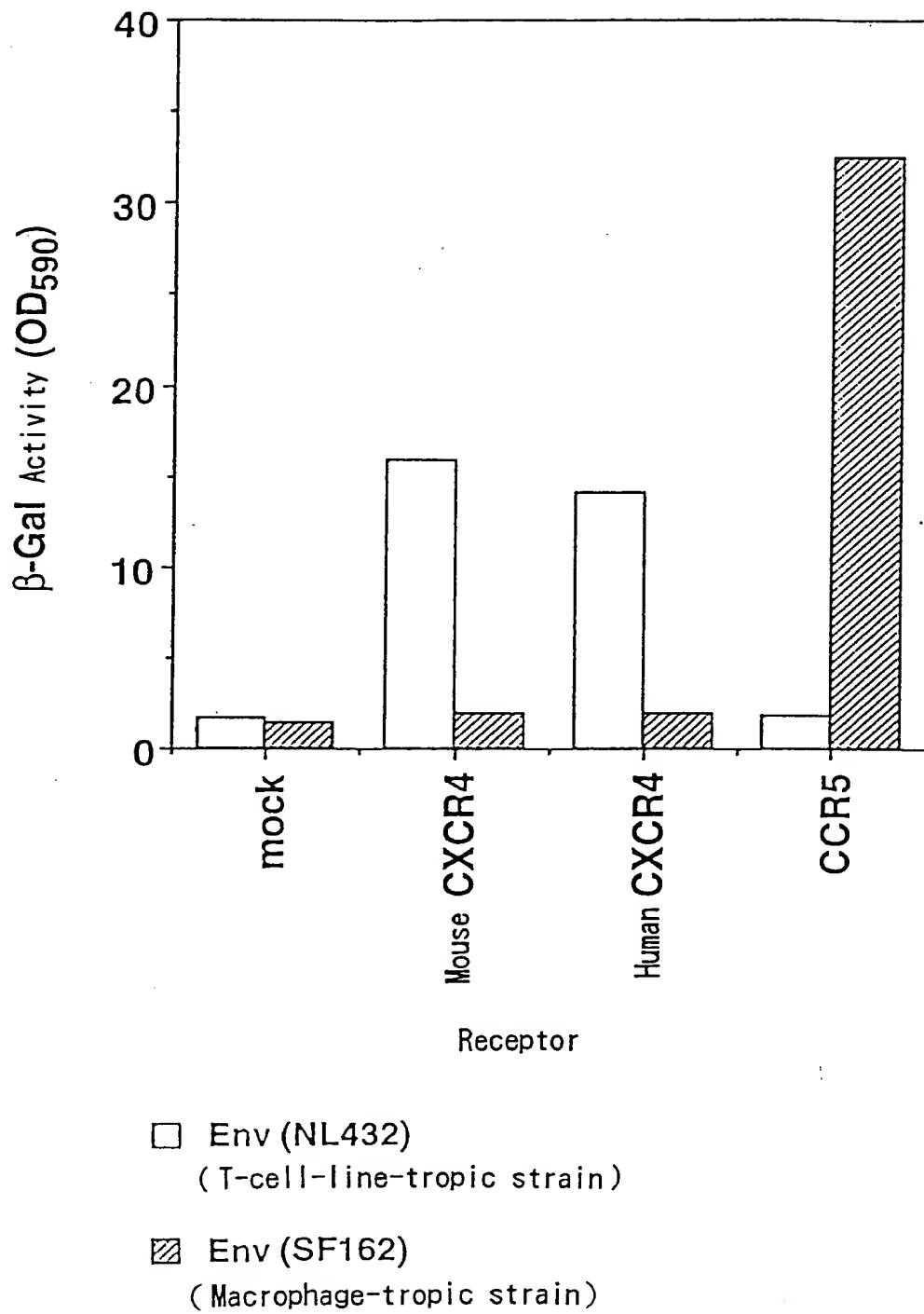


Fig. 8

A

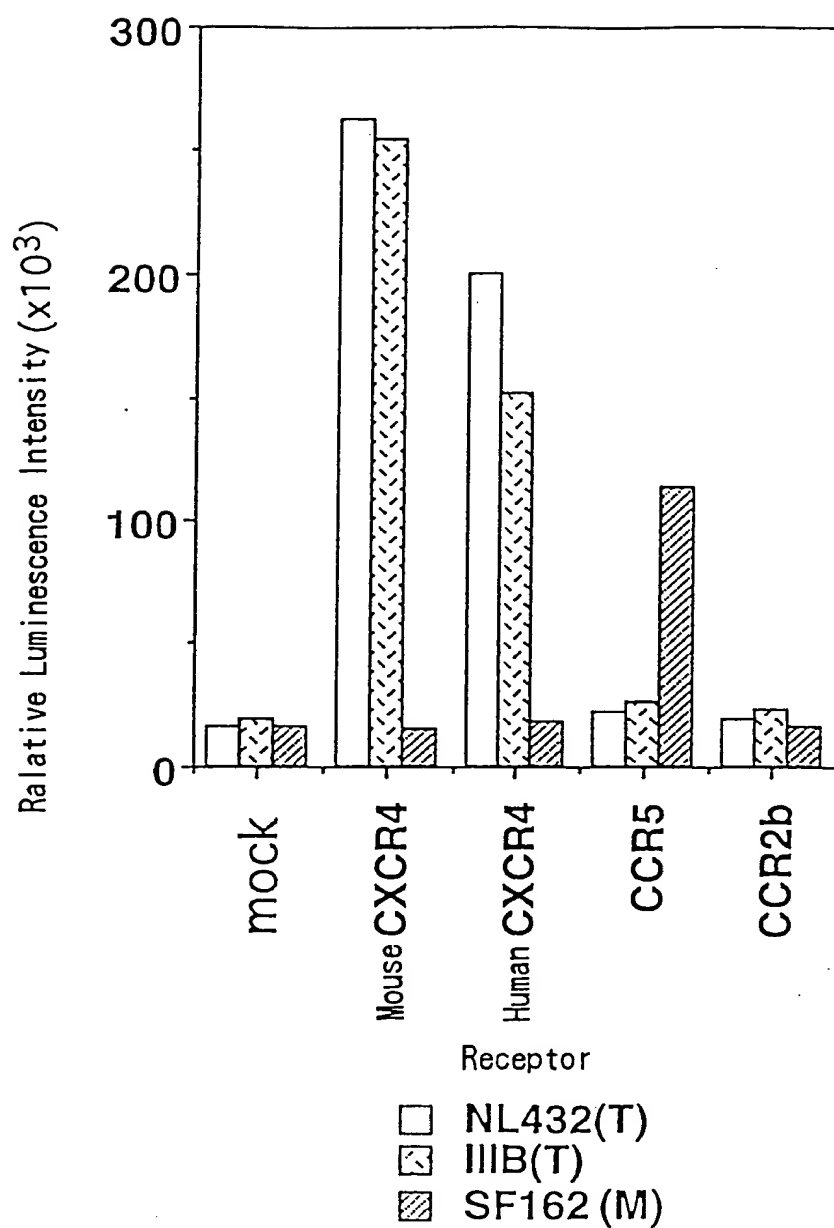


Fig. 9

B

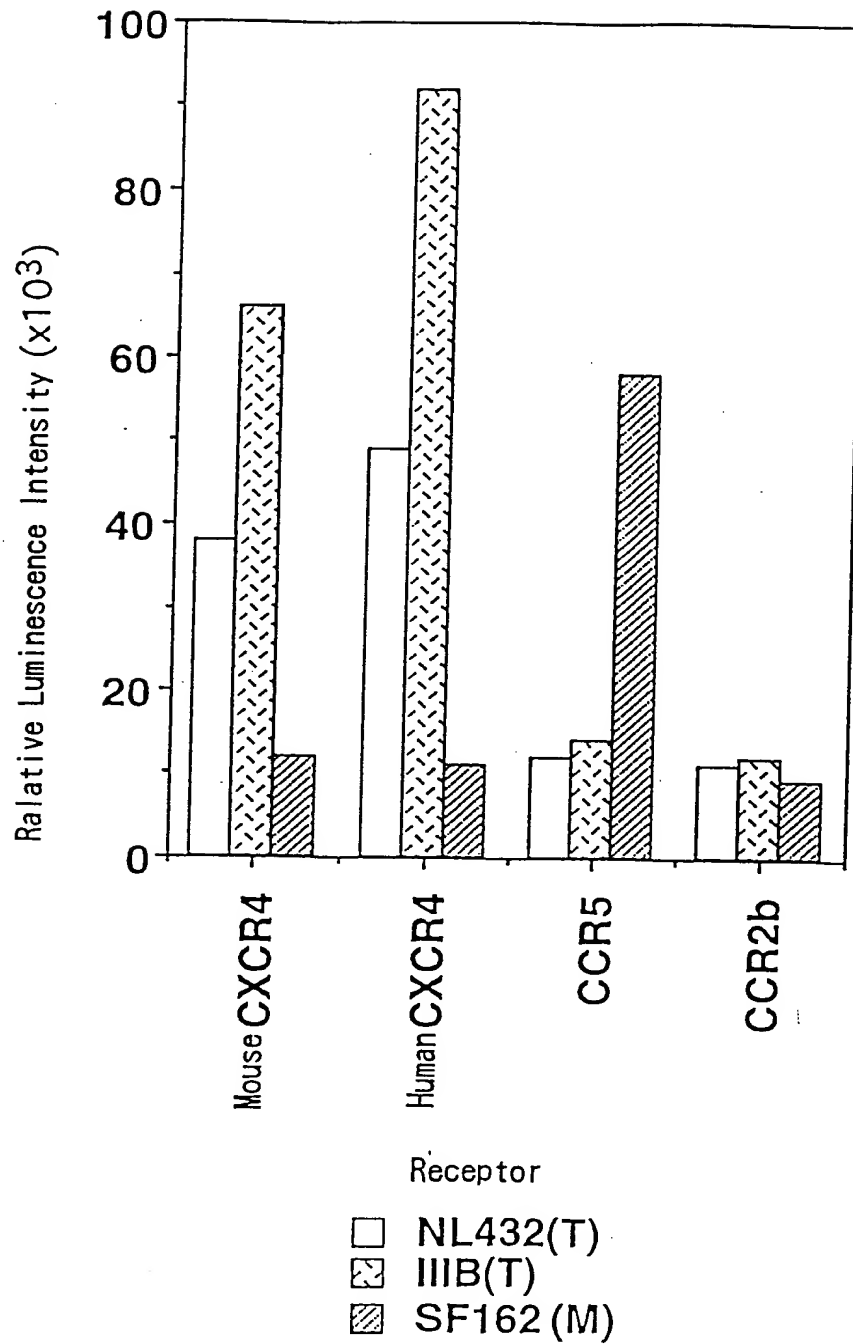


Fig. 10

C

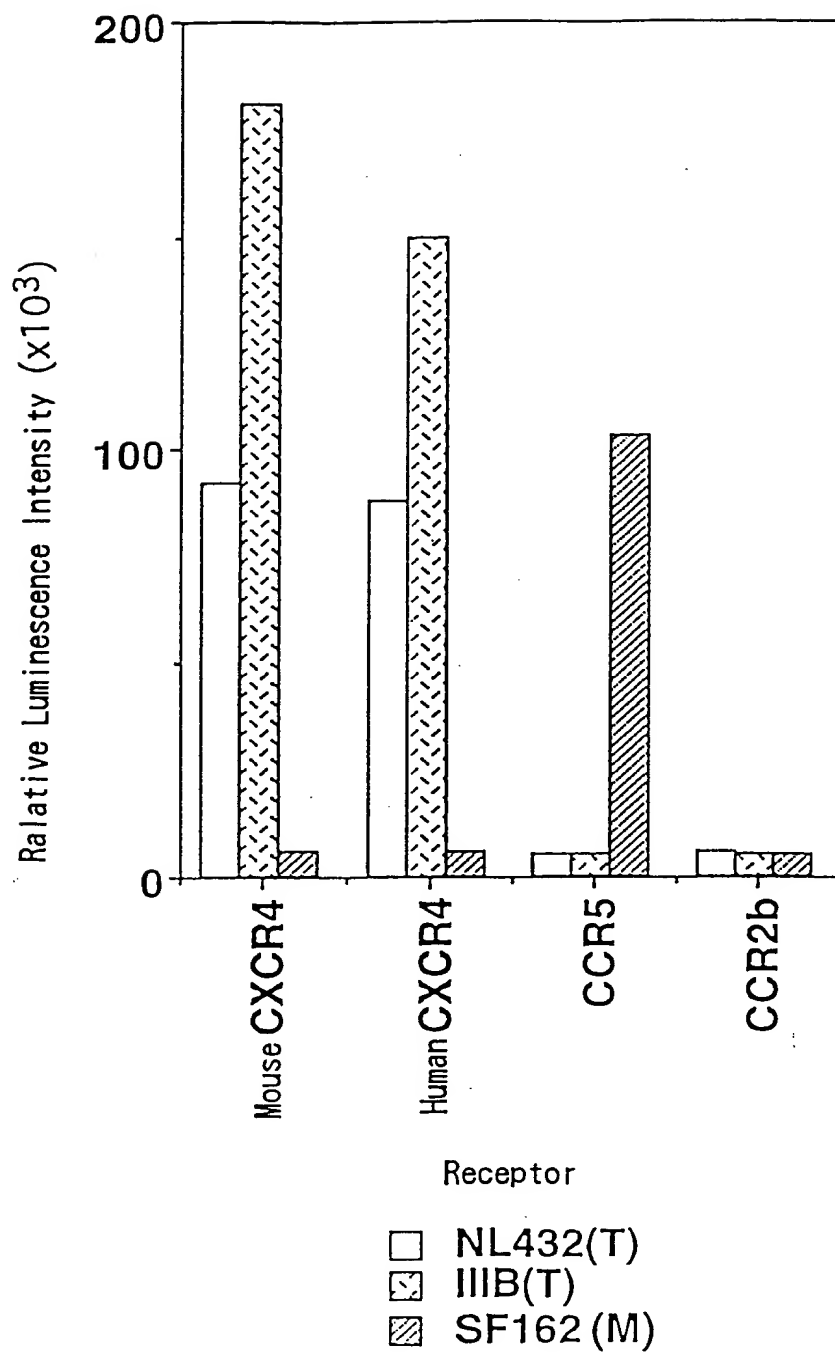


Fig. 11

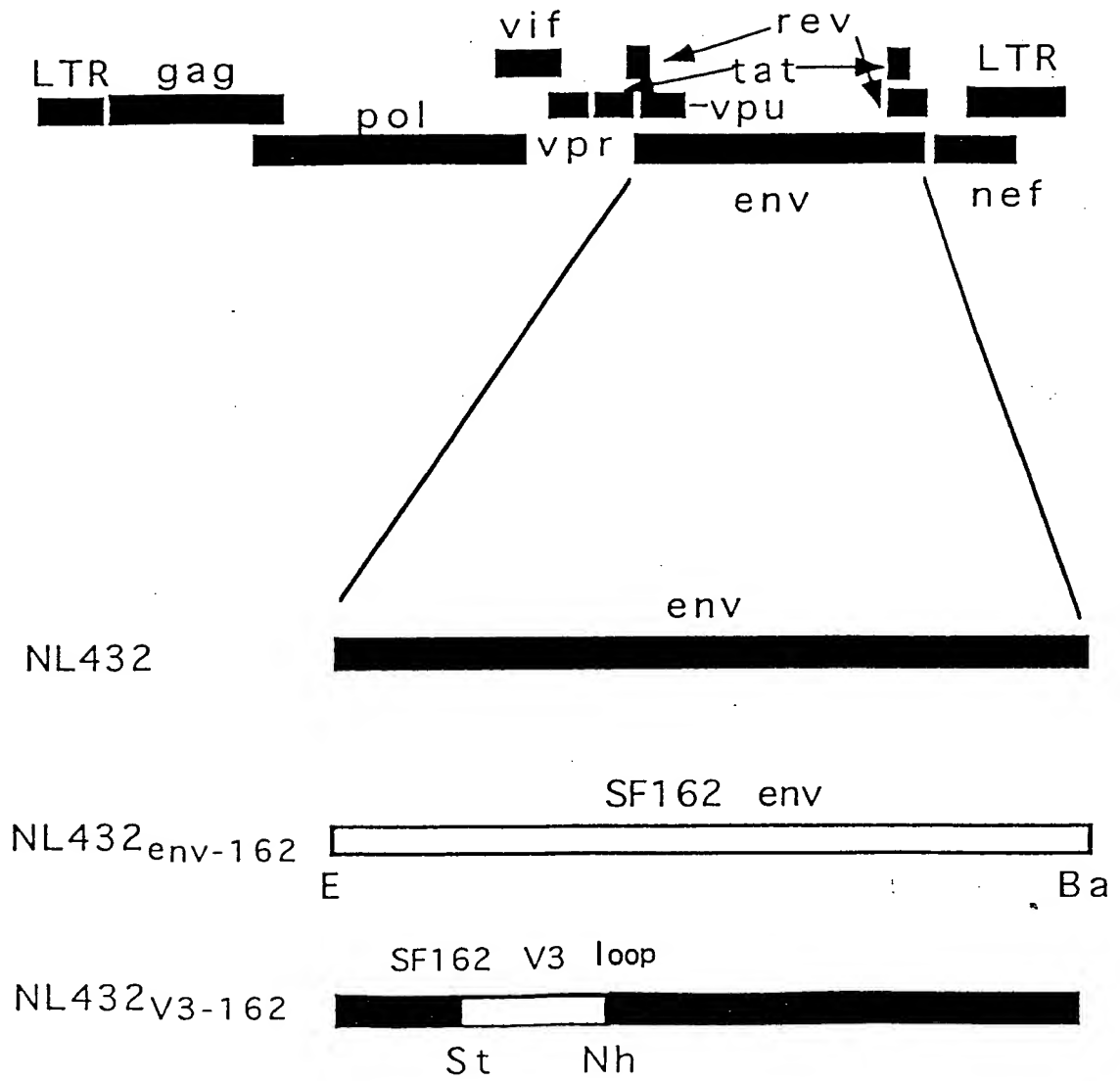


Fig. 12

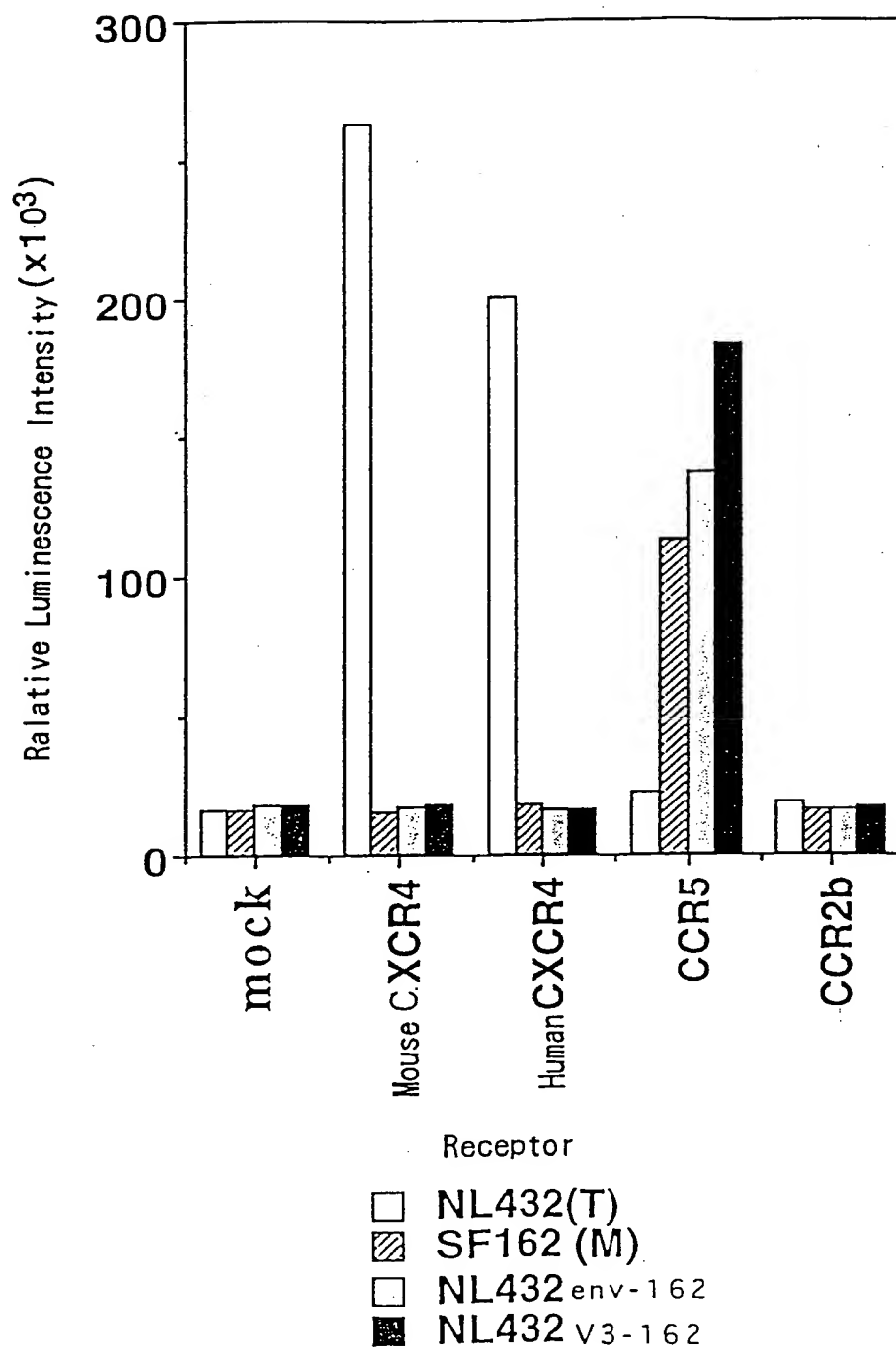


Fig. 13